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Antiviral activity *in vitro* of double combinations of enteroviral inhibitors

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The challenges in combating enteroviruses continue due to their genetic diversity and the rapid development of resistance. Combining antiviral agents offers a promising solution. Our study evaluated the combined effects of pocapavir, a potent capsid-binding inhibitor, with enterovirus-targeting compounds such as pleconaril, 2-(3,4-dichlorophenoxy)-5-nitrobenzonitrile (MDL-860), guanidine hydrochloride, oxoglaucine, and $2-\alpha$ -hydroxybenzylbenzimidazole (HBB). The combinations tested demonstrated additive to highly synergistic antiviral effects without observed combined cytotoxicity. These results suggest their potential for further research and clinical trials in combating enterovirus infections.

KEYWORDS

pocapavir, double combination, CV-B4, CV-A9, PV-1

Introduction

Enteroviruses (EVs) are a group of single, positive-stranded RNA viruses of the *Picornaviridae* family that include polioviruses, echoviruses, coxsackieviruses and enteroviruses. EVs are associated with several human diseases and affect millions of people of all ages worldwide each year. Clinical manifestations of enteroviral infection differ by viral type. Commonly, infections caused by non-polio enteroviruses are asymptomatic or result only in an undifferentiated febrile illness, sometimes they are associated with severe diseases such as aseptic meningitis, encephalitis, non-polio flaccid paralysis, dilated cardiomyopathy (Rotbart, 2002). Treatment for these infections is mainly supportive. The combat against enteroviruses follows two main directions: the development and implementation of effective vaccines and the discovery of chemotherapeutic agents that selectively inhibit viral replication.

The historical Salk and Sabin vaccines against polio are at present the only ones applied in clinical practice. To date several vaccines against enterovirus 71 are available or under development in China (Li et al., 2014; Liang and Wang, 2014). Candidate vaccines containing B4, B5, C1, C2, and C4 genogroups are in development (Chong et al., 2014; Chia et al., 2018; Tamura et al., 2022). Vaccines against Coxsackie A16, A6 and A10 is being considered with a view to developing combination/multivalent vaccines (Lim et al., 2018; He et al., 2021; Zhang et al., 2023). Other vaccines such as recombinant and other

forms of subunit vaccines, vectored vaccines, virus-like particle vaccines are in the early stage of development (Zhang et al., 2010; He et al., 2021).

The fast development of drug resistance is the main problem in the chemotherapy of enteroviral infections, based until now on monotherapy courses (Loddo, 1980). Several capsid-binding agents (such as pleconaril, disoxaril, pirodavir, vapendavir) as well as 3C protease inhibitors (such as rupintrivir), 3A protein Inhibitors (such as enviroxime) failed in several clinical trials of human enterovirus infection due to limited efficacy or side effects (Tammaro et al., 2023). Currently, pocapavir is undergoing Phase2a, and like pleconaril it is available for use in emergency treatment of severe EV-B infections. Thus, there is a need for more potent therapy, as well as for treatment that may decrease the frequency of the emergence of drug-resistant viruses. One way to fight resistance is by applying drug combination therapy of new or already existing substances with different mechanisms of action.

The main aims of using drug combinations are to achieve a synergistic therapeutic effect, reduce dose and toxicity reduction, and to minimize or delay the induction of drug resistance (Chou, 2006). In a synergistic drug combination, the effect is larger than the additive effect of each individual drug, while antagonistic combination has smaller than the additive effect of each individual drug. An experimental alternative strategy based on the synergistic dual combinations of antivirals against the replication of enteroviruses has been established in previous work (Eggers and Tamm, 1963a; Nikolaeva and Galabov, 1999; Nikolaeva and Galabov, 2000; Nikolaeva-Glomb et al., 2011; Thibaut et al., 2011; Wang et al., 2016; Stoyanova and Galabov, 2021a; Lanko et al., 2021; Ianevski et al., 2022; Ianevski et al., 2024).

In the present study, we tested the combined effects of dual combinations involving pocapavir and a panel of picornavirus inhibitors, each with known mechanisms of action. Pocapavir, an orally active capsid inhibitor, falls within the picornavirus antiviral mechanistic class known as capsid inhibitors. It functions by blocking virus uncoating and preventing the release of viral RNA into cells, thus inhibiting virus replication (Wittekind et al., 2017). Pleconaril inhibits virus uncoating by direct insertion into the canyon of enteroviral capsid protein VP1 (Pevear et al., 1999). 2-(3,4dichlorophenoxy)-5-nitrobenzonitrile (MDL-860) targets the host phosphatidylinositol-4 kinase III beta (PI4KB) (Arita et al., 2017), while oxoglaucine, an enviroxime-like inhibitor, specifically targets the PI4KB (Arita et al., 2015). Guanidine hydrochloride interferes with the function of the enteroviral 2C protein, thereby preventing the initiation of negative RNA strand synthesis (Loddo et al., 1962; Barton and Flanegan, 1997). 2-ahydroxybenzyl-benzimidazole (HBB) inhibits the activity of the virus-specific RNA-dependent RNA polymerase, resulting in the suppression of single-stranded RNA synthesis (Eggers and Tamm, 1961; Dmitrieva and Agol, 1974).

Materials and methods

Cells and viruses

Human epithelial type 2 (HEp-2) cells (National Bank for Industrial Microorganisms and Cell Cultures, Sofia, Bulgaria) were grown in Dulbecco's Modified Eagle's Medium (DMEM) supplemented with 10% (v/v) Fetal Bovine Serum (FBS) and antibiotics (penicillin: 100 IU/mL; streptomycin: 100 lg/ml). The cells were maintained in a 37°C incubator with 5% CO₂ and high humidity. The viruses used in this study included Poliovirus type 1 (PV-1), Coxsackievirus A9 (CV-A9) and Coxsackievirus B4 (CV-B4) (obtained from the collection of The Stephan Angeloff Institute of Microbiology, Bulgarian Academy of Sciences, Sofia). They were produced in HEp-2 cells in maintenance medium consisting of DMEM, supplemented with 0.5% FBS and antibiotics. All pathogens were handled in the BSL-2 laboratory.

Antiviral compounds

Pocapavir (SCH-48973; V-073) was purchased from MedKoo Biosciences (Morrisville, United States); pleconaril (WIN 63843) was synthesized and kindly provided by Dr. Vadim Makarov (State Research Center for Antibiotics, Moscow, Russia); MDL-860 [2-(3,4-dichlorophenoxy)-5-nitrobenzonitrile] was synthesized by Dr. Gerhard Purstinger (University of Innsbruck, Austria); oxoglaucine (1,2,9,10-tetramethoxy-7h-dibenzo [de,g]quinolin-7-one) an aporphinoid alkaloid from Glaucium flavum Cranz, was provided by Dr. Stefan Philipov (Institute of Organic Chemistry, BAS); guanidine hydrochloride was obtained from Eastman Organic Chemicals (New York, United States); HBB (2-a-hydroxybenzylbenzimidazole) was received by Dr. T. Dmitrieva (Moscow State University, Moscow, Russia). The compounds were prepared as 10 mmol stock solutions dissolved as follows: pleconaril, MDL-860, and HBB in polyethylene glycol (PEG) 400, pocapavir and oxoglaucine in a 1:9 v/v dimethyl sulfoxide (DMSO)/saline solution, guanidine hydrochloride in a saline solution. These stock solutions were then serially diluted to the desired concentrations in DMEM with 0.5% FBS.

Cell culture assays

The assays were performed using 96-well microplates, infected with approximately 100 CCID_{50} per 0.1 mL of virus. The antiviral effects of the compounds were determined by a cytopathic effect (CPE) inhibition assay. After 1 h of virus adsorption, the compounds were added in various concentrations, and cells were incubated for 48 h at 37°C with 5% CO₂ and high humidity. Inhibition of cytopathic effect was determined using a neutral red uptake assay, based on the initial protocol described by Borenfreund and Puerner (1984). The 50% inhibitory concentration (IC₅₀) of the compounds was identified as the concentration that inhibited the

Pocapavir	Pleconaril		MDL-860	Oxoglaucine	HBB		Guanidine.HCl	
					NH NH	H H H		
Combination		PV-1		CV-A	CV-A9		CV-B4	
		Synergy/Antagonism*		Synergy/Ant	Synergy/Antagonism*		Synergy/Antagonism*	
Pocapavir + Pleconaril		-		0/-4	0/-4.5		60.9/0	
Pocapavir + MDL-860		55.9/0		0.1/0	0.1/0		56.2/0	
Pocapavir + Guanidine.HCl		1.3/0		0/-13	0/-13.4		154.7/0	
Pocapavir + Oxoglaucine		87.2/0		53.2/	53.2/0		276.6/-0.1	
Pocapavir + HBB		0/-21.8		0/-12	0/-12.9		0/-0.4	

TABLE 1 Chemical structure of used antivirals and interactions of drug-drug combinations against PV-1, CV-A9 and CV-B4 viruses.

 aMean volumes of synergy/antagonism using the MacSynergy II program based on the 95% confidence level (in $\mu M^2\%$).

development of CPE by 50%. Drug-drug interactions for three replicate assays were analyzed using the three-dimensional model of Prichard and Shipman (1990), using the MacSynergyTM II software program (Prichard et al., 1992) at 95% confidence limits. The program is an analytical tool which describes and quantitates drug interactions, performs all the required calculations necessary to plot dose-response surface, calculate the volume of synergism or antagonism produced and statically evaluates the interactions for significance. The cytotoxicity of each compound and each combination was assessed in parallel by using the same assay without the addition of the virus.

Statistical analysis

Drug-drug interactions were analyzed by the three dimensional method of Prichard and Shipman (1990), and the analysis was conducted with the MacSynergyTM II (version 1.0) software program. MacSynergyTM II graphically plots three dimension interactions that fall above or below a neutral surface (baseline). Interpretation of drug-drug interactions is based on a volume of synergy or antagonism at 95% confidence interval and calculations, which can be positive (synergy), negative (antagonistic), or neutral (no or minimal interactions, additive). Descriptions of additive, synergistic, and antagonistic interactions in this computer model are based on the Bliss independent model. The synergy/

antagonism volumes (μ M²%) of drug combinations at 95% confidence level were defined as follows: values between -25 and +25 indicate additivity; values between ±25 and ±50 indicate minor but significant antagonism or synergy, respectively; values between ±50 and ±100 are interpreted as moderate synergism or antagonism, respectively; values greater than +100 or less than -100 indicate strong synergy or antagonism, respectively; values over ±1,000 are considered to be the result of experimental error or occasional phenomena and are statistically insignificant (Prichard et al., 1992; Prichard et al., 1993).

Results

Initial, a 50% inhibitory concentrations (IC₅₀) of each compound were determined from the dose-response curves obtained by the cytopathic effect (CPE) inhibition assay. The range of obtained mean values of IC₅₀ for each compound alone against each tested virus is as follows: pocapavir 0.09–0.5 μ M, pleconaril 10–25 μ M, MDL-860 0.2–4 μ M, oxoglaucine 0.05–0.2 μ M, guanidine.HCl 350–590 μ M and HBB 25–125 μ M.

As the next step, we assessed the antiviral activities of various compound pairs, each containing pocapavir, in a checkerboard format using a CPE assay. To determine whether the antiviral activities of the drug combinations were synergistic, antagonistic, or additive, we analyzed the experimental data with the MacSynergyTM II program. This program calculated the



theoretical additive interactions based on the Bliss independence theory and represented each dose combination synergy and antagonism as peaks above or below a predicted additive plane (baseline) in a three-dimensional graph. The corresponding mean volumes of synergy and antagonism at the 95% confidence level are summarized in Table 1.

Effect of double combinations against PV-1 virus

The data for the effect of double combinations against PV-1 virus are presented in Table 1 and Figure 1. When pocapavir was individually combined with MDL-860, or oxoglaucine, it resulted in high volumes of synergy (55.9 and 87.2 μ M²%, respectively), indicating a moderate synergistic effect. When combined with guanidine.HCl, an additive effect was observed. The combination of pocapavir and HBB showed an additive interaction with increased but not significantly antagonistic (–21.7 μ M²%) results. Pleconaril did not exhibit anti-PV-1 activity at doses under 100 μ M and was therefore not included in the combination.

Effect of double combinations against CV-A9 virus

Figure 2 present the data obtained for drug-drug combinations against CV-A9 virus replication. In contrast to the combination of pocapavir and oxoglaucine, which yielded a moderate synergistic effect (53.2 μ M²%), the other combinations of pocapavir with pleconaril, MDL-860, guanidine.HCl, or HBB resulted in an additive effect with no significant antagonism.

Effect of double combinations against CV-B4 virus

The combinations of pocapavir and oxoglaucine or guanidine.HCl against CV-B4 showed strong synergy (276.6 and 154.7 μ M²%, respectively) and a low level of antagonism (Figure 3). The combinations of pocapavir and pleconaril or MDL-860 showed moderate synergy (60.9 and 56.2 μ M²%, respectively) with no antagonism present.



In summary, these experiments revealed a range of moderate to strong synergistic interactions, especially in the combination of pocapavir and oxoglaucine against all included viruses, particularly in CV-B4-induced CPE inhibition. The degrees of synergistic interactions between pocapavir and pleconaril or MDL-860 were essentially the same for PV-1 and CV-A9 inhibition based on the volume of synergy values. Interactions between pocapavir and guanidine.HCl were strong synergistic for CV-B4 CPE inhibition, and additive for PV-1 and CV-A9. The combined effects of pocapavir and HBB were additive.



All combination studies included evaluations of cell viability, and none of the synergistic and additive combinations showed cytotoxicity within the range of drug concentrations examined (Figure 4).

Discussion

Enteroviruses are widespread and economically significant, making them a significant challenge for researchers. Their



combinations using the MacSynergy II program. Data shown were obtained at the 95% confidence level. Values in the zero plane indicate additive activity, values under the zero plane indicate antagonistic activity, and values above the zero plane indicate synergistic activity. All data points are averages of four measurements from at least three independent experiments.

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genetic diversity and rapid development of resistance make them particularly hard to combat. To address this, our team is exploring various methods, such as the use of drug combinations, which can reduce the likelihood of drugresistant mutants emerging (Stoyanova and Galabov, 2020; Stoyanova and Galabov, 2021a; Stoyanova and Galabov, 2021b). The use of drug combinations reduces the possibility of selecting drug-resistant mutants (Mouton, 1999). Additionally, drug combinations that leverage synergistic interactions can enhance antiviral efficacy while minimizing cytotoxicity by reducing the required therapeutic doses. Combining antiviral agents with different resistance profiles and modes of action creates a higher genetic barrier to resistance development, suppressing its emergence.

Our study aimed to evaluate the combined effects of pocapavir and compounds with anti-enterovirus activity and a known mode of action against different enteroviruses. Pocapavir is a potent capsidbinding inhibitor that blocks virus uncoating and viral RNA release, thus preventing virus replication. It had previously shown promise in in vitro/vivo activities and had undergone phase II clinical trials for poliomyelitis treatment, but no recent reports on its development were identified (Collett et al., 2017). In this study, we selected five other compounds based on their mechanisms of action for combination studies and conducted two-drug combination experiments. Pleconaril, like pocapavir, inhibits virus uncoating by binding to the viral particle in the pocket of enteroviral structural protein VP1 (Pevear et al., 1999), which stabilizes the capsid, prevents conformational changes, and inhibits viral genome release. The rapid development of resistance to capsid binders is a major problem in the clinical development of this class of compounds. MDL-860 is a broad-spectrum antipicornavirus compound, it targets the host PI4KB and causes covalent modification and irreversible inactivation of the PI4KB (Arita et al., 2017). Oxoglaucine is an inhibitor of the PI4KB/oxysterol-binding protein (OSBP) pathway in EV replication, directly targeting the PI4KB (Arita et al., 2015). Guanidine hydrochloride and HBB suggest similar but not identical mechanisms of antiviral action, by blocking viral RNA synthesis (Eggers and Tamm, 1963b). Guanidine hydrochloride interferes with the function of the 2C protein, one of the most conserved polyproteins of the EVs, thereby preventing the initiation of negative RNA strand synthesis (Loddo et al., 1962; Barton and Flanegan, 1997). HBB blocks viral RNA synthesis by inhibiting the activity of the enteroviral RNA-dependent RNA polymerase (Eggers and Tamm, 1961; Dmitrieva and Agol, 1974).

The combinations of pocapavir and any of the five other compounds produced at least additive antiviral effects on virusinduced CPE inhibition. This suggests that pocapavir's mode of action is not affected by the mechanisms of the other compounds, based on the absence of significant antagonism. The highest synergistic effect for each of the used viruses was observe with the combination of pocapavir and oxoglaucine. A similar, albeit more modest, result was observed in the combination of pocapavir and MDL-860. Oxoglaucine and MDL-860 have similar modes of antiviral action. The combined effect of pocapavir and MDL-860 was moderately synergistic against PV-1 and CV-B4, while for CV-A9, the combined effect was additive. Based on these results, we can speculate that, the combination of capsid-binding inhibitor and compounds blocking host proteins, providing key stages of viral RNA replication, has a moderate to strong inhibitory effect on viral progeny, depending on the virus. On the other hand, in this case, combining an entry inhibitor with inhibitors of a specific viral RNA synthesis, does not result in an increased inhibitory effect, except in combination with guanidine hydrochloride against CV-A9. The combined effect of pocapavir and HBB was additive against each of the three viruses, while the effect of combination with guanidine hydrochloride resulted in an additive effect against PV-1 and CV-A9, and a synergistic effect against CV-B4. Similar results were obtained in previous studies involving other capsid-binding inhibitors (Nikolaeva and Galabov, 1999; Nikolaeva-Glomb and Galabov, 2004; Stoyanova and Galabov, 2021a). Pocapavir combined with oxoglaucine demonstrated a synergistic effect, while the combination of pleconaril or disoxaril and oxoglaucine had an additive effect. The effect of combination of pocapavir and pleconaril resulted in moderate synergism against CV-B4 replication, and an additive effect on CV-A9. Based on the obtained results, it is noticed that CV-B4 has a higher sensitivity to the examined combinations.

The differences in observed effects among individual virus strains can be attributed to variations in sensitivity to tested compounds. This suggests that since the mechanism of action of the tested compounds is known and enteroviruses share a similar replicative cycle, differences in combined effects may stem from individual variances in target region and thus the compound's impact on specific viral strains. However, additional studies are needed to clarify this statement.

In summary, four of the analyzed two-drug combinations (i.e., pocapavir plus pleconaril, pocapavir plus MDL-860, pocapavir plus oxoglaucine and pocapavir plus guanidine.HCl) demonstrate moderate to strong synergistic antiviral effects, without cytotoxicity, in HEp-2 cell culture. These results suggest the potential for further *in vivo* research and provide a basis for combining anti-enteroviral agents in clinical trials to combat enteroviral infections.

Data availability statement

The raw data supporting the conclusion of this article will be made available by the authors, without undue reservation.

Ethics statement

Ethical approval was not required for the studies on humans in accordance with the local legislation and institutional requirements because only commercially available established cell lines were used.

Author contributions

Conceptualization, AS and AG; methodology, AS and SG; formal analysis, AS and SG; investigation, AS and SG; data curation, AS; writing—original draft preparation, AS, AG, and SG; project administration, AS; funding acquisition, AS. All authors contributed to the article and approved the submitted version.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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